

08135602.TXT  
SEQUENCE LISTING

<110> SOEJIMA, KENJI  
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MAEDA, HIROAKI  
NOZAKI, CHIKATERU  
HAMAMOTO, TAKAYOSHI  
NAKAGAKI, TOMOHIRO

<120> ANTIBODY AGAINST VON WILLEBRAND FACTOR CLEAVING ENZYME  
AND ASSAY SYSTEM USING THE SAME

<130> 081356-0237

<140> 10/529,009

<141> 2005-03-24

<150> PCT/JP03/12280

<151> 2003-09-25

<150> JP 2002-279924

<151> 2002-09-25

<150> JP 2002-377569

<151> 2002-12-26

<160> 24

<170> PatentIn version 3.5

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<212> PRT

<213> Homo sapiens

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Phe Gln Gln Ser Cys Leu Gln Ala Leu Glu Pro Gln Ala Val Ser Ser  
35 40 45

Tyr Leu Ser Pro Gly Ala Pro Leu Lys Gly Arg Pro Pro Ser Pro Gly  
50 55 60

Phe Gln Arg Gln Arg Gln Arg Gln Arg Arg Ala Ala Gly Gly Ile Leu  
65 70 75 80

His Leu Glu Leu Leu Val Ala Val Gly Pro Asp Val Phe Gln Ala His  
85 90 95

Gln Glu Asp Thr Glu Arg Tyr Val Leu Thr Asn Leu Asn Ile Gly Ala  
100 105 110

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Glu Leu Leu Arg Asp Pro Ser Leu Gly Ala Gln Phe Arg Val His Leu  
 115 120 125  
 Val Lys Met Val Ile Leu Thr Glu Pro Glu Gly Ala Pro Asn Ile Thr  
 130 135 140  
 Ala Asn Leu Thr Ser Ser Leu Leu Ser Val Cys Gly Trp Ser Gln Thr  
 145 150 155 160  
 Ile Asn Pro Glu Asp Asp Thr Asp Pro Gly His Ala Asp Leu Val Leu  
 165 170 175  
 Tyr Ile Thr Arg Phe Asp Leu Glu Leu Pro Asp Gly Asn Arg Gln Val  
 180 185 190  
 Arg Gly Val Thr Gln Leu Gly Gly Ala Cys Ser Pro Thr Trp Ser Cys  
 195 200 205  
 Leu Ile Thr Glu Asp Thr Gly Phe Asp Leu Gly Val Thr Ile Ala His  
 210 215 220  
 Glu Ile Gly His Ser Phe Gly Leu Glu His Asp Gly Ala Pro Gly Ser  
 225 230 235 240  
 Gly Cys Gly Pro Ser Gly His Val Met Ala Ser Asp Gly Ala Ala Pro  
 245 250 255  
 Arg Ala Gly Leu Ala Trp Ser Pro Cys Ser Arg Arg Gln Leu Leu Ser  
 260 265 270  
 Leu Leu Ser Ala Gly Arg Ala Arg Cys Val Trp Asp Pro Pro Arg Pro  
 275 280 285  
 Gln Pro Gly Ser Ala Gly His Pro Pro Asp Ala Gln Pro Gly Leu Tyr  
 290 295 300  
 Tyr Ser Ala Asn Glu Gln Cys Arg Val Ala Phe Gly Pro Lys Ala Val  
 305 310 315 320  
 Ala Cys Thr Phe Ala Arg Glu His Leu Asp Met Cys Gln Ala Leu Ser  
 325 330 335  
 Cys His Thr Asp Pro Leu Asp Gln Ser Ser Cys Ser Arg Leu Leu Val  
 340 345 350  
 Pro Leu Leu Asp Gly Thr Glu Cys Gly Val Glu Lys Trp Cys Ser Lys  
 355 360 365

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Gly Arg Cys Arg Ser Leu Val Glu Leu Thr Pro Ile Ala Ala Val His  
370 375 380

Gly Arg Trp Ser Ser Trp Gly Pro Arg Ser Pro Cys Ser Arg Ser Cys  
385 390 395 400

Gly Gly Gly Val Val Thr Arg Arg Arg Gln Cys Asn Asn Pro Arg Pro  
405 410 415

Ala Phe Gly Gly Arg Ala Cys Val Gly Ala Asp Leu Gln Ala Glu Met  
420 425 430

Cys Asn Thr Gln Ala Cys Glu Lys Thr Gln Leu Glu Phe Met Ser Gln  
435 440 445

Gln Cys Ala Arg Thr Asp Gly Gln Pro Leu Arg Ser Ser Pro Gly Gly  
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Ala Ser Phe Tyr His Trp Gly Ala Ala Val Pro His Ser Gln Gly Asp  
465 470 475 480

Ala Leu Cys Arg His Met Cys Arg Ala Ile Gly Glu Ser Phe Ile Met  
485 490 495

Lys Arg Gly Asp Ser Phe Leu Asp Gly Thr Arg Cys Met Pro Ser Gly  
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Pro Arg Glu Asp Gly Thr Leu Ser Leu Cys Val Ser Gly Ser Cys Arg  
515 520 525

Thr Phe Gly Cys Asp Gly Arg Met Asp Ser Gln Gln Val Trp Asp Arg  
530 535 540

Cys Gln Val Cys Gly Gly Asp Asn Ser Thr Cys Ser Pro Arg Lys Gly  
545 550 555 560

Ser Phe Thr Ala Gly Arg Ala Arg Glu Tyr Val Thr Phe Leu Thr Val  
565 570 575

Thr Pro Asn Leu Thr Ser Val Tyr Ile Ala Asn His Arg Pro Leu Phe  
580 585 590

Thr His Leu Ala Val Arg Ile Gly Gly Arg Tyr Val Val Ala Gly Lys  
595 600 605

Met Ser Ile Ser Pro Asn Thr Thr Tyr Pro Ser Leu Leu Glu Asp Gly

610

615

Arg 625	Val	Glu	Tyr	Arg	Val 630	Ala	Leu	Thr	Glu	Asp 635	Arg	Leu	Pro	Arg	Leu 640
Glu	Glu	Ile	Arg	Ile 645	Trp	Gly	Pro	Leu	Gln 650	Glu	Asp	Ala	Asp	Ile 655	Gln
Val	Tyr	Arg	Arg 660	Tyr	Gly	Glu	Glu	Tyr 665	Gly	Asn	Leu	Thr	Arg 670	Pro	Asp
Ile	Thr	Phe 675	Thr	Tyr	Phe	Gln	Pro 680	Lys	Pro	Arg	Gln	Ala 685	Trp	Val	Trp
Ala	Ala 690	Val	Arg	Gly	Pro	Cys 695	Ser	Val	Ser	Cys	Gly 700	Ala	Gly	Leu	Arg
Trp 705	Val	Asn	Tyr	Ser	Cys 710	Leu	Asp	Gln	Ala	Arg 715	Lys	Glu	Leu	Val	Glu 720
Thr	Val	Gln	Cys	Gln 725	Gly	Ser	Gln	Gln	Pro 730	Pro	Ala	Trp	Pro	Glu 735	Ala
Cys	Val	Leu	Glu 740	Pro	Cys	Pro	Pro	Tyr 745	Trp	Ala	Val	Gly	Asp 750	Phe	Gly
Pro	Cys	Ser 755	Ala	Ser	Cys	Gly	Gly 760	Gly	Leu	Arg	Glu	Arg 765	Pro	Val	Arg
Cys	Val 770	Glu	Ala	Gln	Gly	Ser 775	Leu	Leu	Lys	Thr	Leu 780	Pro	Pro	Ala	Arg
Cys 785	Arg	Ala	Gly	Ala	Gln 790	Gln	Pro	Ala	Val	Ala 795	Leu	Glu	Thr	Cys	Asn 800
Pro	Gln	Pro	Cys	Pro 805	Ala	Arg	Trp	Glu	Val 810	Ser	Glu	Pro	Ser	Ser 815	Cys
Thr	Ser	Ala	Gly 820	Gly	Ala	Gly	Leu	Ala 825	Leu	Glu	Asn	Glu	Thr 830	Cys	Val
Pro	Gly	Ala 835	Asp	Gly	Leu	Glu	Ala 840	Pro	Val	Thr	Glu	Gly 845	Pro	Gly	Ser
Val	Asp 850	Glu	Lys	Leu	Pro	Ala 855	Pro	Glu	Pro	Cys	Val 860	Gly	Met	Ser	Cys

Pro 865 Pro Gly Trp Gly 870 His Leu Asp Ala Thr 875 Ser Ala Gly Glu Lys Ala 880

Pro Ser Pro Trp Gly 885 Ser Ile Arg Thr 890 Gly Ala Gln Ala Ala His Val 895

Trp Thr Pro 900 Ala Ala Gly Ser Cys 905 Ser Val Ser Cys Gly Arg Gly Leu 910

Met Glu 915 Leu Arg Phe Leu Cys 920 Met Asp Ser Ala Leu Arg 925 Val Pro Val

Gln 930 Glu Glu Leu Cys Gly 935 Leu Ala Ser Lys Pro Gly 940 Ser Arg Arg Glu

Val 945 Cys Gln Ala Val 950 Pro Cys Pro Ala Arg 955 Trp Gln Tyr Lys Leu Ala 960

Ala Cys Ser Val 965 Ser Cys Gly Arg Gly 970 Val Val Arg Arg Ile Leu Tyr 975

Cys Ala Arg 980 Ala His Gly Glu Asp 985 Asp Gly Glu Glu Ile Leu Leu Asp 990

Thr Gln 995 Cys Gln Gly Leu Pro Arg 1000 Pro Glu Pro Gln Glu 1005 Ala Cys Ser

Leu 1010 Glu Pro Cys Pro Pro Arg 1015 Trp Lys Val Met 1020 Ser Leu Gly Pro

Cys 1025 Ser Ala Ser Cys Gly 1030 Leu Gly Thr Ala Arg 1035 Arg Ser Val Ala

Cys 1040 Val Gln Leu Asp Gln Gly 1045 Gln Asp Val Glu 1050 Val Asp Glu Ala

Ala 1055 Cys Ala Ala Leu Val Arg 1060 Pro Glu Ala Ser 1065 Val Pro Cys Leu

Ile 1070 Ala Asp Cys Thr Tyr Arg 1075 Trp His Val Gly 1080 Thr Trp Met Glu

Cys 1085 Ser Val Ser Cys Gly 1090 Asp Gly Ile Gln Arg 1095 Arg Arg Asp Thr

Cys 1100 Leu Gly Pro Gln Ala Gln 1105 Ala Pro Val Pro 1110 Ala Asp Phe Cys

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Pro	Cys	Val	Gly	Gln	Gly	Thr	Pro	Ser	Leu	Val	Pro	His	Glu	Glu
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Ala	Ala	Ala	Pro	Gly	Arg	Thr	Thr	Ala	Thr	Pro	Ala	Gly	Ala	Ser
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Leu	Glu	Trp	Ser	Gln	Ala	Arg	Gly	Leu	Leu	Phe	Ser	Pro	Ala	Pro
	1160					1165					1170			
Gln	Pro	Arg	Arg	Leu	Leu	Pro	Gly	Pro	Gln	Glu	Asn	Ser	Val	Gln
	1175					1180					1185			
Ser	Ser	Ala	Cys	Gly	Arg	Gln	His	Leu	Glu	Pro	Thr	Gly	Thr	Ile
	1190					1195					1200			
Asp	Met	Arg	Gly	Pro	Gly	Gln	Ala	Asp	Cys	Ala	Val	Ala	Ile	Gly
	1205					1210					1215			
Arg	Pro	Leu	Gly	Glu	Val	Val	Thr	Leu	Arg	Val	Leu	Glu	Ser	Ser
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Leu	Asn	Cys	Ser	Ala	Gly	Asp	Met	Leu	Leu	Leu	Trp	Gly	Arg	Leu
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Thr	Trp	Arg	Lys	Met	Cys	Arg	Lys	Leu	Leu	Asp	Met	Thr	Phe	Ser
	1250					1255					1260			
Ser	Lys	Thr	Asn	Thr	Leu	Val	Val	Arg	Gln	Arg	Cys	Gly	Arg	Pro
	1265					1270					1275			
Gly	Gly	Gly	Val	Leu	Leu	Arg	Tyr	Gly	Ser	Gln	Leu	Ala	Pro	Glu
	1280					1285					1290			
Thr	Phe	Tyr	Arg	Glu	Cys	Asp	Met	Gln	Leu	Phe	Gly	Pro	Trp	Gly
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Glu	Ile	Val	Ser	Pro	Ser	Leu	Ser	Pro	Ala	Thr	Ser	Asn	Ala	Gly
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Gly	Cys	Arg	Leu	Phe	Ile	Asn	Val	Ala	Pro	His	Ala	Arg	Ile	Ala
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Ile	His	Ala	Leu	Ala	Thr	Asn	Met	Gly	Ala	Gly	Thr	Glu	Gly	Ala
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Asn Ala Ser Tyr Ile Leu Ile Arg Asp Thr His Ser Leu Arg Thr  
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Thr Ala Phe His Gly Gln Gln Val Leu Tyr Trp Glu Ser Glu Ser  
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Ser Gln Ala Glu Met Glu Phe Ser Glu Gly Phe Leu Lys Ala Gln  
1385 1390 1395

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tt 62

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